

(Z)-4-Bromo-N-[(Z)-3-[(4-bromo-2,6-diisopropylphenyl)imino]butan-2-ylidene]-2,6-diisopropylaniline

Cun Zhang, Guo-Fan Wu, Bao-Mei Huang and Xiao-Quan Lu*

Key Laboratory of Bioelectrochemistry & Environmental Analysis of Gansu Province, College of Chemistry & Chemical Engineering, Northwest Normal University, Lanzhou 730070, People's Republic of China

Correspondence e-mail: luxq@nwnu.edu.cn

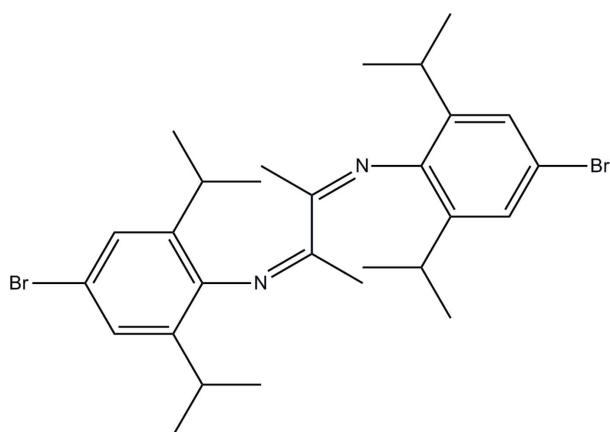
Received 11 December 2012; accepted 27 December 2012

Key indicators: single-crystal X-ray study; $T = 296$ K; mean $\sigma(\text{C}-\text{C}) = 0.009$ Å; R factor = 0.052; wR factor = 0.179; data-to-parameter ratio = 17.9.

The title compound, $\text{C}_{28}\text{H}_{38}\text{Br}_2\text{N}_2$, is centrosymmetric with the mid-point of the central C–C bond of the butyl group located on an inversion center. The terminal benzene ring is approximately perpendicular to the central 1,4-diaza-butadiene mean plane [dihedral angle = 78.23 (3)°]. No hydrogen bonding or aromatic stacking is observed in the crystal structure.

Related literature

For applications of diimine catalysts, see: Cotts *et al.* (2000); Ittel *et al.* (2000); Johnson *et al.* (1995); Zhang & Ye (2012).



Experimental

Crystal data

$\text{C}_{28}\text{H}_{38}\text{Br}_2\text{N}_2$
 $M_r = 562.42$
Monoclinic, $P2_1/n$
 $a = 9.099$ (3) Å
 $b = 12.199$ (4) Å
 $c = 13.566$ (5) Å
 $\beta = 104.905$ (5)°

$V = 1455.2$ (9) Å³
 $Z = 2$
Mo $K\alpha$ radiation
 $\mu = 2.80$ mm⁻¹
 $T = 296$ K
 $0.25 \times 0.23 \times 0.19$ mm

Data collection

Bruker APEXII CCD
diffractometer
Absorption correction: multi-scan
(SADABS; Bruker, 2001)
 $T_{\min} = 0.541$, $T_{\max} = 0.618$

7266 measured reflections
2685 independent reflections
1460 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.043$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.052$
 $wR(F^2) = 0.179$
 $S = 0.94$
2685 reflections
150 parameters

84 restraints
H-atom parameters constrained
 $\Delta\rho_{\max} = 0.46$ e Å⁻³
 $\Delta\rho_{\min} = -0.40$ e Å⁻³

Data collection: *APEX2* (Bruker, 2007); cell refinement: *SAINT* (Bruker, 2007); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL* (Sheldrick, 2008); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

This work was supported by the National Natural Science Foundation of China (Nos. 20927004, 21175108 and 20875077), the Natural Science Foundation of Gansu Province (Nos. 0803RJZA105, 096RJZA121 and 096RJZA122) and the Key Laboratory of Polymer Materials of Gansu Province, China.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: XU5664).

References

Bruker (2001). *SADABS*. Bruker AXS Inc., Madison, Wisconsin, USA.
Bruker (2007). *APEX2* and *SAINT*. Bruker AXS Inc., Madison, Wisconsin, USA.
Cotts, P. M., Guan, Z., McCord, E. F. & McLain, S. J. (2000). *Macromolecules*, **33**, 6945–6952.
Ittel, S. D., Johnson, L. K. & Brookhart, M. (2000). *Chem. Rev.* **100**, 1169–1203.
Johnson, L. K., Killian, C. M. & Brookhart, M. (1995). *J. Am. Chem. Soc.* **117**, 6414–6415.
Sheldrick, G. M. (2008). *Acta Cryst. A* **64**, 112–122.
Zhang, Z. & Ye, Z. (2012). *Chem. Commun.* **48**, 7940–7942.

supplementary materials

Acta Cryst. (2013). **E69**, o216 [doi:10.1107/S160053681205194X]

(Z)-4-Bromo-N-{(Z)-3-[(4-bromo-2,6-diisopropylphenyl)imino]butan-2-yl- idene}-2,6-diisopropylaniline

Cun Zhang, Guo-Fan Wu, Bao-Mei Huang and Xiao-Quan Lu

Comment

In recent years, Ni(II)/Pd(II)- α -diimine catalysts were greatly attracted attention due to their high catalytic activity and influence on product performance in olefin polymerization (Zhang & Ye, 2012; Johnson *et al.*, 1995). It is well known that the polymerization conditions (such as the reaction olefin pressure, temperature *etc.*) and ligand structure had a great impact on catalytic activity and polymer properties (Cotts *et al.*, 2000; Ittel *et al.*, 2000).

In the solid state, the structure exhibits *trans*-conformation about the central C—C bond of the ligand backbone. Bond lengths and angles are within the expected range for α -diimines. The dihedral angle between the aryl ring and 1,4-diaza-butadiene plane is 78.23 (3) $^{\circ}$ (Fig. 1). In the crystal packing, there is no hydrogen-bond between the molecules.

Experimental

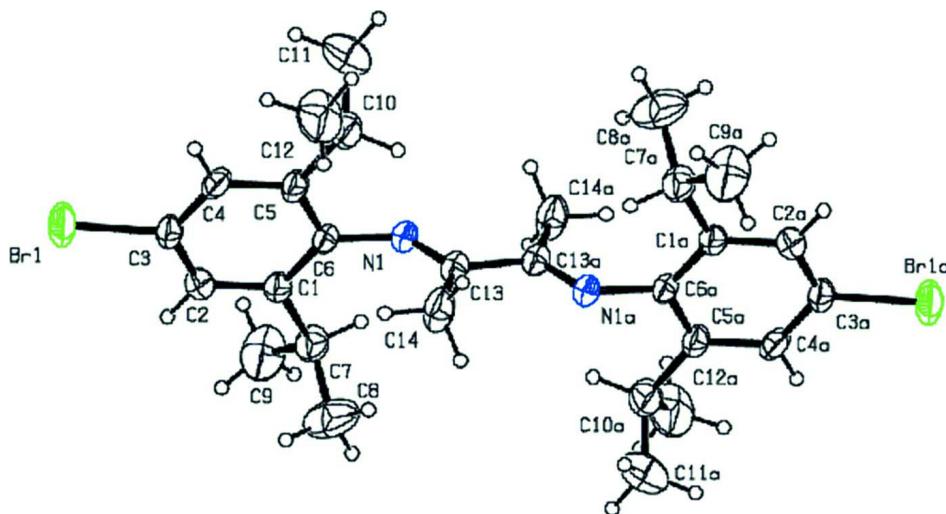
Formic acid (0.5 ml) was added to a stirred solution of 2,3-butanedione (0.042 g, 0.49 mmol) and 4-Bromo-2,6-diisopropyl-phenylamine (0.250 g, 0.98 mmol) in methanol (20 ml). The mixture was refluxed for 24 h, then cooled and the precipitate was separated by filtration. The solid was recrystallized from dichloromethane/cyclohexane (*v/v* = 8:1), washed with cold ethanol and dried under vacuum to give the title ligand 0.21 g (75%). Anal. Calcd. for C₂₈H₃₈Br₂N₂: C, 59.79; H, 6.81; N, 4.98; Found: C, 60.29; H, 6.95; N, 4.74.

Refinement

All hydrogen atoms were placed in calculated positions with C—H distances of 0.93 and 0.96 Å for aryl and methyl type H-atoms. They were included in the refinement in a riding model approximation, respectively. The H-atoms were assigned *U*_{iso} = 1.2 times *U*_{eq} of the aryl C atoms and 1.5 times *U*_{eq} of the methyl C atoms.

Computing details

Data collection: *APEX2* (Bruker, 2007); cell refinement: *SAINT* (Bruker, 2007); data reduction: *SAINT* (Bruker, 2007); program(s) used to solve structure: *SHELXTL* (Sheldrick, 2008); program(s) used to refine structure: *SHELXTL* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL* (Sheldrick, 2008).

**Figure 1**

Molecular structure of the title compound, using 30% probability level ellipsoids.

(Z)-4-Bromo-N-[(Z)-3-[(4-bromo-2,6-diisopropylphenyl)imino]butan-2-ylidene]-2,6-diisopropylaniline

Crystal data



$M_r = 562.42$

Monoclinic, $P2_1/n$

Hall symbol: -P 2yn

$a = 9.099 (3) \text{ \AA}$

$b = 12.199 (4) \text{ \AA}$

$c = 13.566 (5) \text{ \AA}$

$\beta = 104.905 (5)^\circ$

$V = 1455.2 (9) \text{ \AA}^3$

$Z = 2$

$F(000) = 580$

$D_x = 1.284 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$

Cell parameters from 1370 reflections

$\theta = 2.9\text{--}21.4^\circ$

$\mu = 2.80 \text{ mm}^{-1}$

$T = 296 \text{ K}$

Block, yellow

$0.25 \times 0.23 \times 0.19 \text{ mm}$

Data collection

Bruker APEXII CCD
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

φ and ω scans

Absorption correction: multi-scan
(*SADABS*; Bruker, 2001)

$T_{\min} = 0.541$, $T_{\max} = 0.618$

7266 measured reflections

2685 independent reflections

1460 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.043$

$\theta_{\max} = 25.5^\circ$, $\theta_{\min} = 2.3^\circ$

$h = -10 \rightarrow 10$

$k = -12 \rightarrow 14$

$l = -16 \rightarrow 14$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.052$

$wR(F^2) = 0.179$

$S = 0.94$

2685 reflections

150 parameters

84 restraints

Primary atom site location: structure-invariant
direct methods

Secondary atom site location: difference Fourier

map

Hydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.1P)^2 + 0.7029P]$
where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\text{max}} < 0.001$

$\Delta\rho_{\max} = 0.46 \text{ e \AA}^{-3}$

$\Delta\rho_{\min} = -0.40 \text{ e \AA}^{-3}$

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) etc. and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
Br1	0.35688 (8)	0.65673 (6)	0.10127 (7)	0.0922 (4)
C1	0.7029 (6)	0.8851 (4)	0.1392 (4)	0.0489 (12)
C2	0.5759 (6)	0.8270 (4)	0.1476 (4)	0.0533 (14)
H2	0.5247	0.8477	0.1958	0.064*
C3	0.5254 (6)	0.7392 (5)	0.0851 (4)	0.0539 (14)
C4	0.5971 (7)	0.7079 (4)	0.0117 (4)	0.0557 (15)
H4	0.5608	0.6482	-0.0301	0.067*
C5	0.7232 (7)	0.7651 (4)	-0.0001 (5)	0.0585 (14)
C6	0.7756 (6)	0.8543 (4)	0.0646 (4)	0.0452 (13)
C7	0.7611 (8)	0.9814 (5)	0.2083 (5)	0.0662 (14)
H7	0.8701	0.9872	0.2130	0.079*
C8	0.6907 (11)	1.0865 (6)	0.1624 (7)	0.126 (3)
H8A	0.7162	1.0991	0.0990	0.188*
H8B	0.7283	1.1459	0.2084	0.188*
H8C	0.5822	1.0819	0.1505	0.188*
C9	0.7469 (11)	0.9636 (8)	0.3149 (6)	0.120 (3)
H9A	0.8150	1.0122	0.3607	0.180*
H9B	0.7725	0.8891	0.3348	0.180*
H9C	0.6443	0.9783	0.3175	0.180*
C10	0.8059 (9)	0.7265 (6)	-0.0773 (6)	0.0804 (16)
H10	0.8666	0.7876	-0.0923	0.096*
C11	0.9126 (12)	0.6347 (7)	-0.0315 (8)	0.132 (3)
H11A	0.9847	0.6606	0.0287	0.199*
H11B	0.9656	0.6101	-0.0802	0.199*
H11C	0.8555	0.5750	-0.0139	0.199*
C12	0.6986 (12)	0.6892 (8)	-0.1768 (7)	0.126 (3)
H12A	0.6336	0.6323	-0.1632	0.189*
H12B	0.7566	0.6616	-0.2214	0.189*
H12C	0.6378	0.7500	-0.2086	0.189*
C13	0.9228 (6)	0.9765 (4)	-0.0034 (4)	0.0512 (14)
C14	0.7932 (7)	1.0159 (6)	-0.0879 (5)	0.081 (2)
H14A	0.8104	0.9956	-0.1523	0.122*
H14B	0.7856	1.0942	-0.0843	0.122*
H14C	0.7003	0.9831	-0.0812	0.122*
N1	0.9141 (5)	0.9060 (3)	0.0633 (3)	0.0504 (11)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Br1	0.0812 (6)	0.0846 (6)	0.1286 (8)	-0.0381 (4)	0.0595 (5)	-0.0188 (4)
C1	0.053 (3)	0.045 (3)	0.054 (3)	-0.005 (2)	0.023 (2)	-0.001 (2)
C2	0.053 (3)	0.054 (3)	0.058 (4)	-0.001 (3)	0.023 (3)	-0.001 (3)
C3	0.055 (3)	0.052 (3)	0.060 (4)	-0.011 (3)	0.026 (3)	0.005 (3)
C4	0.070 (4)	0.034 (3)	0.066 (4)	-0.014 (3)	0.024 (3)	-0.004 (3)
C5	0.075 (3)	0.043 (3)	0.070 (3)	-0.012 (3)	0.043 (3)	0.001 (3)
C6	0.046 (3)	0.037 (3)	0.055 (3)	-0.005 (2)	0.017 (3)	0.008 (2)
C7	0.076 (3)	0.062 (3)	0.067 (3)	-0.015 (3)	0.028 (3)	-0.015 (3)
C8	0.145 (7)	0.070 (5)	0.145 (6)	-0.001 (5)	0.005 (6)	-0.036 (5)
C9	0.155 (7)	0.131 (6)	0.083 (5)	-0.052 (5)	0.045 (5)	-0.038 (5)
C10	0.100 (4)	0.066 (3)	0.095 (4)	-0.016 (3)	0.060 (3)	-0.018 (3)
C11	0.131 (7)	0.129 (6)	0.163 (7)	0.022 (5)	0.085 (6)	-0.026 (6)
C12	0.153 (7)	0.147 (6)	0.100 (6)	-0.030 (6)	0.073 (5)	-0.040 (5)
C13	0.053 (3)	0.045 (3)	0.059 (4)	-0.006 (2)	0.019 (3)	0.007 (3)
C14	0.059 (4)	0.083 (5)	0.093 (5)	-0.012 (3)	0.005 (4)	0.039 (4)
N1	0.052 (3)	0.044 (3)	0.059 (3)	-0.006 (2)	0.022 (2)	0.005 (2)

Geometric parameters (\AA , $^\circ$)

Br1—C3	1.894 (5)	C9—H9A	0.9600
C1—C2	1.384 (7)	C9—H9B	0.9600
C1—C6	1.396 (7)	C9—H9C	0.9600
C1—C7	1.511 (8)	C10—C11	1.507 (11)
C2—C3	1.371 (7)	C10—C12	1.517 (11)
C2—H2	0.9300	C10—H10	0.9800
C3—C4	1.376 (8)	C11—H11A	0.9600
C4—C5	1.387 (7)	C11—H11B	0.9600
C4—H4	0.9300	C11—H11C	0.9600
C5—C6	1.402 (7)	C12—H12A	0.9600
C5—C10	1.513 (8)	C12—H12B	0.9600
C6—N1	1.414 (7)	C12—H12C	0.9600
C7—C8	1.495 (10)	C13—N1	1.265 (6)
C7—C9	1.501 (9)	C13—C14	1.497 (8)
C7—H7	0.9800	C13—C13 ⁱ	1.498 (10)
C8—H8A	0.9600	C14—H14A	0.9600
C8—H8B	0.9600	C14—H14B	0.9600
C8—H8C	0.9600	C14—H14C	0.9600
C2—C1—C6	118.8 (5)	H9A—C9—H9B	109.5
C2—C1—C7	120.9 (5)	C7—C9—H9C	109.5
C6—C1—C7	120.2 (5)	H9A—C9—H9C	109.5
C3—C2—C1	120.2 (5)	H9B—C9—H9C	109.5
C3—C2—H2	119.9	C11—C10—C5	109.2 (6)
C1—C2—H2	119.9	C11—C10—C12	110.1 (7)
C2—C3—C4	121.2 (5)	C5—C10—C12	112.8 (6)
C2—C3—Br1	119.7 (4)	C11—C10—H10	108.2
C4—C3—Br1	119.0 (4)	C5—C10—H10	108.2

C3—C4—C5	120.3 (5)	C12—C10—H10	108.2
C3—C4—H4	119.8	C10—C11—H11A	109.5
C5—C4—H4	119.8	C10—C11—H11B	109.5
C4—C5—C6	118.4 (5)	H11A—C11—H11B	109.5
C4—C5—C10	119.8 (5)	C10—C11—H11C	109.5
C6—C5—C10	121.7 (5)	H11A—C11—H11C	109.5
C1—C6—C5	121.1 (5)	H11B—C11—H11C	109.5
C1—C6—N1	118.6 (5)	C10—C12—H12A	109.5
C5—C6—N1	119.9 (5)	C10—C12—H12B	109.5
C8—C7—C9	113.0 (7)	H12A—C12—H12B	109.5
C8—C7—C1	111.3 (5)	C10—C12—H12C	109.5
C9—C7—C1	112.5 (5)	H12A—C12—H12C	109.5
C8—C7—H7	106.5	H12B—C12—H12C	109.5
C9—C7—H7	106.5	N1—C13—C14	125.7 (5)
C1—C7—H7	106.5	N1—C13—C13 ⁱ	116.6 (6)
C7—C8—H8A	109.5	C14—C13—C13 ⁱ	117.8 (6)
C7—C8—H8B	109.5	C13—C14—H14A	109.5
H8A—C8—H8B	109.5	C13—C14—H14B	109.5
C7—C8—H8C	109.5	H14A—C14—H14B	109.5
H8A—C8—H8C	109.5	C13—C14—H14C	109.5
H8B—C8—H8C	109.5	H14A—C14—H14C	109.5
C7—C9—H9A	109.5	H14B—C14—H14C	109.5
C7—C9—H9B	109.5	C13—N1—C6	122.2 (4)
C6—C1—C2—C3	1.6 (8)	C4—C5—C6—N1	172.3 (5)
C7—C1—C2—C3	−179.3 (5)	C10—C5—C6—N1	−4.1 (9)
C1—C2—C3—C4	−1.3 (9)	C2—C1—C7—C8	−90.6 (8)
C1—C2—C3—Br1	177.3 (4)	C6—C1—C7—C8	88.5 (8)
C2—C3—C4—C5	0.2 (9)	C2—C1—C7—C9	37.4 (8)
Br1—C3—C4—C5	−178.4 (4)	C6—C1—C7—C9	−143.5 (6)
C3—C4—C5—C6	0.4 (9)	C4—C5—C10—C11	−81.6 (8)
C3—C4—C5—C10	176.9 (6)	C6—C5—C10—C11	94.7 (8)
C2—C1—C6—C5	−0.9 (8)	C4—C5—C10—C12	41.1 (9)
C7—C1—C6—C5	180.0 (5)	C6—C5—C10—C12	−142.6 (7)
C2—C1—C6—N1	−173.4 (5)	C14—C13—N1—C6	0.5 (9)
C7—C1—C6—N1	7.5 (8)	C13 ⁱ —C13—N1—C6	−179.2 (6)
C4—C5—C6—C1	−0.1 (8)	C1—C6—N1—C13	−106.2 (6)
C10—C5—C6—C1	−176.4 (6)	C5—C6—N1—C13	81.2 (7)

Symmetry code: (i) $-x+2, -y+2, -z$.